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SEQ ID NO:1. Human Hepsin DNA Sequence: The GenBank Accession No. for human hepsin mRNA (cDNA) is M18930:

	1	TCGAGCCCGC	TTTCCAGGGA	CCCTACCTGA	GGGCCCACAG	GTGAGGCAGC	CTGGCCTAGC
5	61	AGGCCCCACG	CCACCGCCTC	TGCCTCCAGG	CCGCCCGCTG	CTGCGGGGCC	ACCATGCTCC
J	121	TGCCCAGGCC	TGGAGACTGA	CCCGACCCCG	GCACTACCTC	GAGGCTCCGC	CCCCACCTGC
	181	TGGACCCCAG	GGTCCCACCC	TGGCCCAGGA	GGTCAGCCAG	GGAATCATTA	ACAAGAGGCA
	241	GTGACATGGC	GCAGAAGGAG	GGTGGCCGGA	CTGTGCCATG	CTGCTCCAGA	CCCAAGGTGG
	301	CAGCTCTCAC	TGCGGGGACC	CTGCTACTTC	TGACAGCCAT	CGGGGCGCA	TCCTGGGCCA
10	361	ттстссстст	TCTCCTCAGG	AGTGACCAGG	AGCCGCTGTA	CCCAGTGCAG	GTCAGCTCTG
10	421	CGGACGCTCG	GCTCATGGTC	TTTGACAAGA	CGGAAGGGAC	GTGGCGGCTG	CTGTGCTCCT
	481	CGCGCTCCAA	CGCCAGGGTA	GCCGGACTCA	GCTGCGAGGA	GATGGGCTTC	CTCAGGGCAC
	541	TGACCCACTC	CGAGCTGGAC	GTGCGAACGG	CGGGCGCCAA	TGGCACGTCG	GGCTTCTTCT
	601	GTGTGGACGA	GGGGAGGCTG	CCCCACACCC	AGAGGCTGCT	GGAGGTCATC	TCCGTGTGTG
15	661	ATTGCCCCAG	AGGCCGTTTC	TTGGCCGCCA	TCTGCCAAGA	CTGTGGCCGC	
10	721		CATCGTGGGA			CCGGTGGCCG	
	781	GCCTTCGCTA	TGATGGAGCA	CACCTCTGTG	GGGGATCCCT		GACTGGGTGC
	841	TGACAGCCGC	CCACTGCTTC	CCGGAGCGGA	ACCGGGTCCT	GTCCCGATGG	
	901	CCGGTGCCGT	GGCCCAGGCC	TCTCCCCACG	GTCTGCAGCT		GCTGTGGTCT
20	961	ACCACGGGGG	CTATCTTCCC	TTTCGGGACC	CCAACAGCGA	GGAGAACAGC	AACGATATTG
	1021		CCTCTCCAGT		TCACAGAATA		
	1081	CAGCTGCCGG	CCAGGCCCTG	GTGGATGGCA	AGATCTGTAC	0010/100000	
	1141	CGCAGTACTA	TGGCCAACAG	GCCGGGGTAC	TCCAGGAGGC		ATAATCAGCA
	1201	ATGATGTCTG	CAATGGCGCT		GAAACCAGAT		
25	1261	CTGGCTACCC	CGAGGGTGGC	ATTGATGCCT	GCCAGGGCGA	CAGCGGTGGT	CCCTTTGTGT
	1321	GTGAGGACAG				TGGCATTGTG	AGTTGGGGCA
	1381	CTGGCTGTGC	CCTGGCCCAG	AAGCCAGGCG			TTCCGGGAGT
	1441	GGATCTTCCA			AAGCCAGCGG	CATGGTGACC	CAGCTCTGAC
	1501	CGGTGGCTTC		GCCTCCAGGG			GGGATCCACG
30	1561	CTGGGCCGAG	GATGGGACGT	TTTTCTTCTT			AGGACACCCT
	1621	CCCTCCAGGG			GGCCCACTCA	GCCCCGAGAC	CACCCAACCT
	1681	CACCCTCCTG	ACCCCCATGT				CTAGGTGCCC
	1741	CTGATGATGG	GATGCTCTTT	AAATAATAAA	GATGGTTTTG	ATT	

SEQ ID NO:2. Human Hepsin Polypeptide Sequence (417 a.a.): The protein_id number is "AAA36013.1:

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NH<sub>2</sub>-

1 MAQKEGGRTV PCCSRPKVAA LTAGTLLLT AIGAASWAIV AVLLRSDQEP LYPVQVSSAD
61 ARLMVFDKTE GTWRLLCSSR SNARVAGLSC EEMGFLRALT HSELDVRTAG ANGTSGFFCV
121 DEGRLPHTQR LLEVISVCDC PRGRFLAAIC QDCGRRKLPV DRIVGGRDTS LGRWPWQVSL
181 RYDGAHLCGG SLLSGDWVLT AAHCFPERNR VLSRWRVFAG AVAQASPHGL QLGVQAVVYH
241 GGYLPFRDPN SEENSNDIAL VHLSSPLPLT EYIQPVCLPA AGQALVDGKI CTVTGWGNTQ
301 YYGQQAGVLQ EARVPIISND VCNGADFYGN QIKPKMFCAG YPEGGIDACQ GDSGGPFVCE
361 DSISRTPRWR LCGIVSWGTG CALAQKPGVY TKVSDFREWI FQAIKTHSEA SGMVTQL
-COOH
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